STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING
ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: _10/549,796_

Source: _PCT/10_

Date Processed by STIC: _10/4/05_

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:
1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY
FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE <u>CHECKER VERSION 4.2.2 PROGRAM</u>, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:
http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.
Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.
Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (**<http://www.uspto.gov/ebc/efs/downloads/documents.htm>** , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05): U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314

Revised.01/24/05

Raw Sequence Listing Error Summary

ERROR DETECTED	SUGGESTED CORRECTION	SERIAL NUMBER: _10/549,796_

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

1____Wrapped Nucleics Wrapped Aminos
The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."

2____Invalid Line Length
The rules require that a line not exceed 72 characters in length. This includes white spaces.

3____Misaligned Amino Numbering
The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers: use space characters, instead.

4____Non-ASCII
The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.

5____Variable Length
Sequence(s)____contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.

6____PatentIn 2.0 "bug"
A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) _____. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.

7____Skipped Sequences (OLD RULES)
Sequence(s) ____ missing. If intentional, please insert the following lines for each skipped sequence:
(2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
(i)   SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
(xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
This sequence is intentionally skipped

Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.

8____Skipped Sequences (NEW RULES)
Sequence(s) ____ missing. If intentional, please insert the following lines for each skipped sequence.
<210> sequence id number
<400> sequence id number
000

9____Use of n's or Xaa's (NEW RULES)
Use of n's and/or Xaa's have been detected in the Sequence Listing.
Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.

10____Invalid <213> Response
Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence

11____Use of <220>
Sequence(s) _____ missing the <220> "Feature" and associated numeric identifiers and responses. Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.
(See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32)   (Sec. 1.823 of Sequence Rules)

12____PatentIn 2.0 "bug"
Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.

13____Misuse of n/Xaa
"n" can only represent a single <u>nucleotide</u>; "Xaa" can only represent a single <u>amino acid</u>

AMC – Biotechnology Systems Branch – 09/09/2003

PCT

RAW SEQUENCE LISTING   DATE: 10/04/2005
PATENT APPLICATION: US/10/549,796   TIME: 10:16:05

Input Set : A:\12987_54.txt
Output Set: N:\CRF4\10042005\J549796.raw

```
 3 <110> APPLICANT: Enobia Pharma Inc.
 4        Guy Boileau
 5        Adriana Karaoglanovic Carmona
 6        Marcelo Campos
 7        Maria Aparecida Juliano
 8        Luiz Juliano
10 <120> TITLE OF INVENTION: PHEX SUBSTRATES AND METHODS USING SAME
11 <130> FILE REFERENCE: 12987.54
13 <140> CURRENT APPLICATION NUMBER: US/10/549,796
14 <141> CURRENT FILING DATE: 2005-09-19
16 <160> NUMBER OF SEQ ID NOS: 32
18 <170> SOFTWARE: PatentIn version 3.2
20 <210> SEQ ID NO: 1
21 <211> LENGTH: 7
22 <212> TYPE: PRT
23 <213> ORGANISM: Artificial
25 <220> FEATURE:
26 <223> OTHER INFORMATION: Peptide
28 <400> SEQUENCE: 1
30 Ala Trp Leu Asp Ser Gly Val
31 1               5
34 <210> SEQ ID NO: 2
35 <211> LENGTH: 4
36 <212> TYPE: PRT
37 <213> ORGANISM: Artificial
39 <220> FEATURE:
40 <223> OTHER INFORMATION: Peptide
42 <400> SEQUENCE: 2
44 Phe Ser Asp Tyr
45 1
48 <210> SEQ ID NO: 3
49 <211> LENGTH: 6
50 <212> TYPE: PRT
51 <213> ORGANISM: Artificial
53 <220> FEATURE:
54 <223> OTHER INFORMATION: peptide
56 <400> SEQUENCE: 3
58 Gly Phe Ser Asp Tyr Lys
59 1               5
62 <210> SEQ ID NO: 4
63 <211> LENGTH: 6
64 <212> TYPE: PRT
65 <213> ORGANISM: Artificial
```

[handwritten annotations: "pg 1-2, 46", "insufficient explanation give source of genetic material (see item 11 on Error Summary Sheet)", "Does Not Comply Corrected Diskette Needed"]

[vertical text right margin: BEST AVAILABLE COPY]

RAW SEQUENCE LISTING   DATE: 10/04/2005
PATENT APPLICATION: US/10/549,796   TIME: 10:16:05

Input Set : A:\12987_54.txt
Output Set: N:\CRF4\10042005\J549796.raw

```
    67 <220> FEATURE:
    68 <223> OTHER INFORMATION: peptide
    71 <220> FEATURE:
    72 <221> NAME/KEY: misc_feature
    73 <222> LOCATION: (2)..(5)
    74 <223> OTHER INFORMATION: Xaa can be any naturally occurring amino acid except cysteine
    77 <400> SEQUENCE: 4
W--> 79 Gly Xaa Xaa Xaa Xaa Lys
    80 1               5
    83 <210> SEQ ID NO: 5
    84 <211> LENGTH: 6
    85 <212> TYPE: PRT
    86 <213> ORGANISM: Artificial
    88 <220> FEATURE:
    89 <223> OTHER INFORMATION: peptide
    92 <220> FEATURE:
    93 <221> NAME/KEY: MISC_FEATURE
    94 <222> LOCATION: (2)..(3)
    95 <223> OTHER INFORMATION: Xaa can be any naturally occurring amino acid except
cysteine,
    96         aspartate and glutamate
    98 <220> FEATURE:
    99 <221> NAME/KEY: MISC_FEATURE
   100 <222> LOCATION: (5)..(5)
   101 <223> OTHER INFORMATION: Xaa can be any naturally occurring amino acid except
cysteine
   103 <400> SEQUENCE: 5
W--> 105 Gly Xaa Xaa Asp Xaa Lys
   106 1               5
   109 <210> SEQ ID NO: 6
   110 <211> LENGTH: 6
   111 <212> TYPE: PRT
   112 <213> ORGANISM: Artificial
   114 <220> FEATURE:
   115 <223> OTHER INFORMATION: peptide
   118 <220> FEATURE:
   119 <221> NAME/KEY: MISC_FEATURE
   120 <222> LOCATION: (2)..(2)
   121 <223> OTHER INFORMATION: Xaa can be any naturally occurring amino acid except
cysteine,
   122         aspartate and glutamate
   124 <220> FEATURE:
   125 <221> NAME/KEY: MISC_FEATURE
   126 <222> LOCATION: (3)..(3)
   127 <223> OTHER INFORMATION: Xaa can be any naturally occurring amino acid except
cysteine
   129 <220> FEATURE:
   130 <221> NAME/KEY: MISC_FEATURE
   131 <222> LOCATION: (5)..(5)
   132 <223> OTHER INFORMATION: Xaa can be any naturally occurring amino acid except
cysteine,
   133         aspartate and glutamate
   135 <400> SEQUENCE: 6
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W--> 137 Gly Xaa Xaa Asp Xaa Lys

```
    138 1                    5
    141 <210> SEQ ID NO: 7
    142 <211> LENGTH: 6
    143 <212> TYPE: PRT
    144 <213> ORGANISM: Artificial
    146 <220> FEATURE:
    147 <223> OTHER INFORMATION: peptide
    150 <220> FEATURE:
    151 <221> NAME/KEY: MISC_FEATURE
    152 <222> LOCATION: (2)..(2)
    153 <223> OTHER INFORMATION: Xaa can be any naturally occurring amino acid except
cysteine
    155 <220> FEATURE:
    156 <221> NAME/KEY: MISC_FEATURE
    157 <222> LOCATION: (3)..(3)
    158 <223> OTHER INFORMATION: Xaa can be any naturally occurring amino acid except
cysteine,
    159        aspartate and glutamate
    161 <220> FEATURE:
    162 <221> NAME/KEY: MISC_FEATURE
    163 <222> LOCATION: (5)..(5)
    164 <223> OTHER INFORMATION: Xaa can be any naturally occurring amino acid except
cysteine,
    165        aspartate and glutamate
    167 <400> SEQUENCE: 7
W--> 169 Gly Xaa Xaa Asp Xaa Lys
    170 1                    5
    173 <210> SEQ ID NO: 8
    174 <211> LENGTH: 4
    175 <212> TYPE: PRT
    176 <213> ORGANISM: Artificial
    178 <220> FEATURE:
    179 <223> OTHER INFORMATION: peptide
    181 <400> SEQUENCE: 8
    183 Asp Arg Arg Leu
    184 1
    187 <210> SEQ ID NO: 9
    188 <211> LENGTH: 4
    189 <212> TYPE: PRT
    190 <213> ORGANISM: Artificial
    192 <220> FEATURE:
    193 <223> OTHER INFORMATION: peptide
    195 <400> SEQUENCE: 9
    197 Arg His Thr Arg
    198 1
    201 <210> SEQ ID NO: 10
    202 <211> LENGTH: 6
    203 <212> TYPE: PRT
    204 <213> ORGANISM: Artificial
    206 <220> FEATURE:
    207 <223> OTHER INFORMATION: peptide
    210 <220> FEATURE:
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Input Set : A:\12987_54.txt
Output Set: N:\CRF4\10042005\J549796.raw

```
      211 <221> NAME/KEY: misc_feature
      212 <222> LOCATION: (2)..(3)
      213 <223> OTHER INFORMATION: Xaa can be any naturally occurring amino acid
      215 <220> FEATURE:
      216 <221> NAME/KEY: misc_feature
      217 <222> LOCATION: (5)..(5)
      218 <223> OTHER INFORMATION: Xaa can be any naturally occurring amino acid
      220 <400> SEQUENCE: 10
W-->  222 Gly Xaa Xaa Asp Xaa Lys
      223 1               5
      226 <210> SEQ ID NO: 11
      227 <211> LENGTH: 6
      228 <212> TYPE: PRT
      229 <213> ORGANISM: Artificial
      231 <220> FEATURE:
      232 <223> OTHER INFORMATION: peptide
      235 <220> FEATURE:
      236 <221> NAME/KEY: misc_feature
      237 <222> LOCATION: (2)..(3)
      238 <223> OTHER INFORMATION: Xaa can be any naturally occurring amino acid
      240 <220> FEATURE:
      241 <221> NAME/KEY: misc_feature
      242 <222> LOCATION: (5)..(5)
      243 <223> OTHER INFORMATION: Xaa can be any naturally occurring amino acid
      245 <400> SEQUENCE: 11
W-->  247 Gly Xaa Xaa Glu Xaa Lys
      248 1               5
      251 <210> SEQ ID NO: 12
      252 <211> LENGTH: 6
      253 <212> TYPE: PRT
      254 <213> ORGANISM: Artificial
      256 <220> FEATURE:
      257 <223> OTHER INFORMATION: peptide
      259 <400> SEQUENCE: 12
      261 Gly Phe Ser Glu Tyr Lys
      262 1               5
      265 <210> SEQ ID NO: 13
      266 <211> LENGTH: 6
      267 <212> TYPE: PRT
      268 <213> ORGANISM: Artificial
      270 <220> FEATURE:
      271 <223> OTHER INFORMATION: peptide
      273 <400> SEQUENCE: 13
      275 Gly Phe Ser Asp Tyr Gln
      276 1               5
      279 <210> SEQ ID NO: 14
      280 <211> LENGTH: 9
      281 <212> TYPE: PRT
      282 <213> ORGANISM: Artificial
```

```
          Input Set : A:\12987_54.txt
          Output Set: N:\CRF4\10042005\J549796.raw

284 <220> FEATURE:
285 <223> OTHER INFORMATION: peptide
287 <400> SEQUENCE: 14
289 Asp His Leu Ser Asp Thr Ser Thr Gln
290 1               5
293 <210> SEQ ID NO: 15
294 <211> LENGTH: 7
295 <212> TYPE: PRT
296 <213> ORGANISM: Artificial
298 <220> FEATURE:
299 <223> OTHER INFORMATION: peptide
301 <400> SEQUENCE: 15
303 Leu Glu Leu Asp Ser Arg Gln
304 1               5
307 <210> SEQ ID NO: 16
308 <211> LENGTH: 8
309 <212> TYPE: PRT
310 <213> ORGANISM: Artificial
312 <220> FEATURE:
313 <223> OTHER INFORMATION: peptide
315 <400> SEQUENCE: 16
317 Ala Trp Leu Asp Ser Gly Val Gln
318 1               5
321 <210> SEQ ID NO: 17
322 <211> LENGTH: 14
323 <212> TYPE: PRT
324 <213> ORGANISM: Artificial
326 <220> FEATURE:
327 <223> OTHER INFORMATION: peptide
329 <400> SEQUENCE: 17
331 Arg Arg His Thr Arg Ser Ala Glu Asp Asp Ser Glu Arg Gln
332 1               5                        10
335 <210> SEQ ID NO: 18
336 <211> LENGTH: 8
337 <212> TYPE: PRT
338 <213> ORGANISM: Artificial
340 <220> FEATURE:
341 <223> OTHER INFORMATION: peptide
343 <400> SEQUENCE: 18
345 Leu Met Met Asp Phe Arg Gly Gln
346 1               5
349 <210> SEQ ID NO: 19
350 <211> LENGTH: 8
351 <212> TYPE: PRT
352 <213> ORGANISM: Artificial
354 <220> FEATURE:
355 <223> OTHER INFORMATION: peptide
357 <400> SEQUENCE: 19
359 Ser Ala Glu Asp Asn Ser Pro Gln
```

FYI

The types of errors shown exist throughout
the Sequence Listing. Please check subsequent
sequences for similar errors.

global err

RAW SEQUENCE LISTING ERROR SUMMARY   DATE: 10/04/2005
PATENT APPLICATION: US/10/549,796   TIME: 10:16:06

Input Set : A:\12987_54.txt
Output Set: N:\CRF4\10042005\J549796.raw

Please Note:
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the
Sequence Listing to ensure that a corresponding explanation is presented in the <220>
to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:4; Xaa Pos. 2,3,4,,5
Seq#:5; Xaa Pos. 2,3,5
Seq#:6; Xaa Pos. 2,3,5
Seq#:7; Xaa Pos. 2,3,5
Seq#:10; Xaa Pos. 2,3,5
Seq#:11; Xaa Pos. 2,3,5

Invalid <213> Response:
Use of "Artificial" only as "<213> Organism" response is incomplete,
per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:1,2,3,4,5,6,7,8,9,10,11,12,13,14,15,16,17,18,19,20,21,22,23,24,25,26,27
Seq#:28,29,30,31,32

VERIFICATION SUMMARY   DATE: 10/04/2005
PATENT APPLICATION: US/10/549,796   TIME: 10:16:06

Input Set : A:\12987_54.txt
Output Set: N:\CRF4\10042005\J549796.raw

L:13 M:270 C: Current Application Number differs, Replaced Current Application Number
L:14 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:79 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:0
L:105 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:0
L:137 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:0
L:169 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:0
L:222 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:0
L:247 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:0